

**CRF Errors Corrected by the STIC System Branch**

*Team 3*  
*4/02/97*

Serial Number: 08/788,800

CRF Processing Date: \_\_\_\_\_  
Edited by: \_\_\_\_\_  
Verified by: \_\_\_\_\_ (STIC staff)

**INTERVIEWED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☒ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☒ other P0987R1
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_

**\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/788,800DATE: 04/02/97  
TIME: 16:27:39

INPUT SET: S16681.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

## SEQUENCE LISTING

## (1) General Information:

(i) APPLICANT: Bednar, Martin M.  
Thomas, G. Roger  
Gross, Cordell E.

(ii) TITLE OF INVENTION: ANTI-CD18 ANTIBODIES IN STROKE

(iii) NUMBER OF SEQUENCES: 15

## (iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Genentech, Inc.  
(B) STREET: 460 Point San Bruno Blvd  
(C) CITY: South San Francisco  
(D) STATE: California  
(E) COUNTRY: USA  
(F) ZIP: 94080

## (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: WinPatin (Genentech)

## (vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:  
(B) FILING DATE: 22-Jan-1997  
(C) CLASSIFICATION:

## (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Lee, Wendy M.  
(B) REGISTRATION NUMBER: 40,378  
(C) REFERENCE/DOCKET NUMBER: P0987r1

## (ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 415/225-1994  
(B) TELEFAX: 415/952-9881  
(C) TELEX: 910/371-7168

## (2) INFORMATION FOR SEQ ID NO:1:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 98 amino acids  
(B) TYPE: Amino Acid

RAW SEQUENCE LISTING  
PATENT APPLICATION US/68/788,800

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47 (D) TOPOLOGY: Linear

48

49 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

50

51 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser  
52 1 5 10 15

53

54 Lys Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys  
55 20 25 30

56

57 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
58 35 40 45

59

60 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser  
61 50 55 60

62

63 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser  
64 65 70 75

65

66 Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser  
67 80 85 90

68

69 Asn Thr Lys Val Asp Lys Arg Val  
70 95 98

71

72 (2) INFORMATION FOR SEQ ID NO:2:

73

74 (i) SEQUENCE CHARACTERISTICS:

75 (A) LENGTH: 98 amino acids

76 (B) TYPE: Amino Acid

77 (D) TOPOLOGY: Linear

78

79 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

80

81 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser  
82 1 5 10 15

83

84 Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys  
85 20 25 30

86

87 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
88 35 40 45

89

90 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser  
91 50 55 60

92

93 Gly Leu Tyr Ser•Leu Ser Ser Val Val Thr Val Pro Ser Ser Asn  
94 65 70 75

95

96 Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser  
97 80 85 90

98

99 Asn Thr Lys Val Asp Lys Thr Val

RAW SEQUENCE LISTING  
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100 95 98  
101  
102 (2) INFORMATION FOR SEQ ID NO:3:  
103  
104 (i) SEQUENCE CHARACTERISTICS:  
105 (A) LENGTH: 98 amino acids  
106 (B) TYPE: Amino Acid  
107 (D) TOPOLOGY: Linear  
108  
109 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:  
110  
111 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser  
112 1 5 10 15  
113  
114 Arg Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys  
115 20 25 30  
116  
117 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
118 35 40 45  
119  
120 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser  
121 50 55 60  
122  
123 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser  
124 65 70 75  
125  
126 Leu Gly Thr Gln Thr Tyr Thr Cys Asn Val Asn His Lys Pro Ser  
127 80 85 90  
128  
129 Asn Thr Lys Val Asp Lys Arg Val  
130 95 98  
131  
132 (2) INFORMATION FOR SEQ ID NO:4:  
133  
134 (i) SEQUENCE CHARACTERISTICS:  
135 (A) LENGTH: 98 amino acids  
136 (B) TYPE: Amino Acid  
137 (D) TOPOLOGY: Linear  
138  
139 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:  
140  
141 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Cys Ser  
142 1 5 10 15  
143  
144 Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys  
145 20 25 30  
146  
147 Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala  
148 35 40 45  
149  
150 Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser  
151 50 55 60  
152

RAW SEQUENCE LISTING  
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153 Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser  
154 65 70 75

155  
156 Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser  
157 80 85 90

158  
159 Asn Thr Lys Val Asp Lys Arg Val  
160 95 98

161  
162 (2) INFORMATION FOR SEQ ID NO:5:

163  
164 (i) SEQUENCE CHARACTERISTICS:  
165 (A) LENGTH: 107 amino acids  
166 (B) TYPE: Amino Acid  
167 (D) TOPOLOGY: Linear

168  
169 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

170  
171 Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp  
172 1 5 10 15

173  
174 Glu Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn  
175 20 25 30

176  
177 Asn Phe Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn  
178 35 40 45

179  
180 Ala Leu Gln Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp  
181 50 55 60

182  
183 Ser Lys Asp Ser Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser  
184 65 70 75

185  
186 Lys Ala Asp Tyr Glu Lys His Lys Val Tyr Ala Cys Glu Val Thr  
187 80 85 90

188  
189 His Gln Gly Leu Ser Ser Pro Val Thr Lys Ser Phe Asn Arg Gly  
190 95 100 105

191  
192 Glu Cys  
193 107

194  
195 (2) INFORMATION FOR SEQ ID NO:6:

196  
197 (i) SEQUENCE CHARACTERISTICS:  
198 (A) LENGTH: 105 amino acids  
199 (B) TYPE: Amino Acid  
200 (D) TOPOLOGY: Linear

201  
202 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

203  
204 Gln Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser  
205 1 5 10 15

RAW SEQUENCE LISTING  
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206  
207 Glu Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser  
208 20 25 30  
209  
210 Asp Phe Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser  
211 35 40 45  
212  
213 Ser Pro Val Lys Ala Gly Val Glu Thr Thr Thr Pro Ser Lys Gln  
214 50 55 60  
215  
216 Ser Asn Asn Lys Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro  
217 65 70 75  
218  
219 Glu Gln Trp Lys Ser His Arg Ser Tyr Ser Cys Gln Val Thr His  
220 80 85 90  
221  
222 Glu Gly Ser Thr Val Glu Lys Thr Val Ala Pro Thr Glu Cys Ser  
223 95 100 105  
224

## (2) INFORMATION FOR SEQ ID NO:7:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 100 amino acids

(B) TYPE: Amino Acid

(D) TOPOLOGY: Linear

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

233  
234 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Pro  
235 1 5 10 15  
236  
237 Lys Asn Ser Ser Met Ile Ser Asn Thr Pro Ala Leu Gly Cys Leu  
238 20 25 30  
239  
240 Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser  
241 35 40 45  
242  
243 Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln  
244 50 55 60  
245  
246 Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro His  
247 65 70 75  
248  
249 Gln Ser Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys  
250 80 85 90  
251  
252 Pro Ser Asn Thr Lys Val Asp Lys Arg Val  
253 95 100  
254

## (2) INFORMATION FOR SEQ ID NO:8:

## (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 11 amino acids

PAGE: 1

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/788,800**

DATE: 04/02/97

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**INPUT SET: S16681.raw**

Line

Error

Original Text